

SPK-10001 AAV-based microRNA mediates non-allele specific reduction of *HTT* mRNA through RNA interference, demonstrating its potential for further preclinical development

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INTRODUCTION

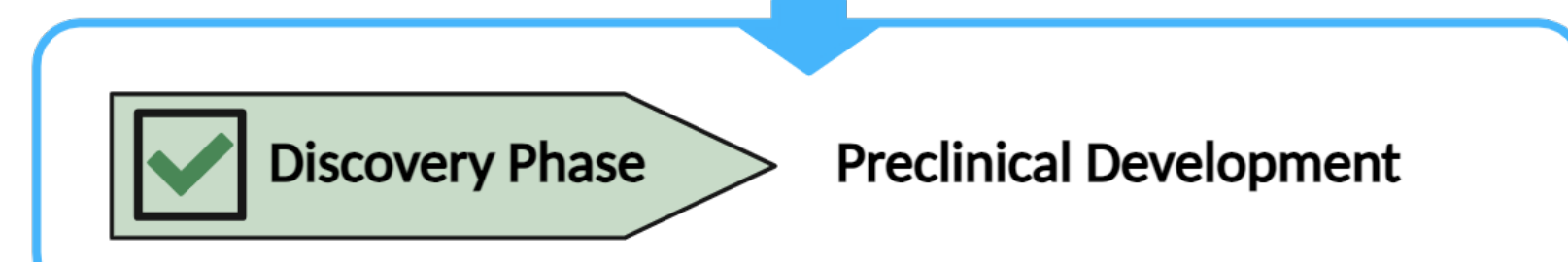
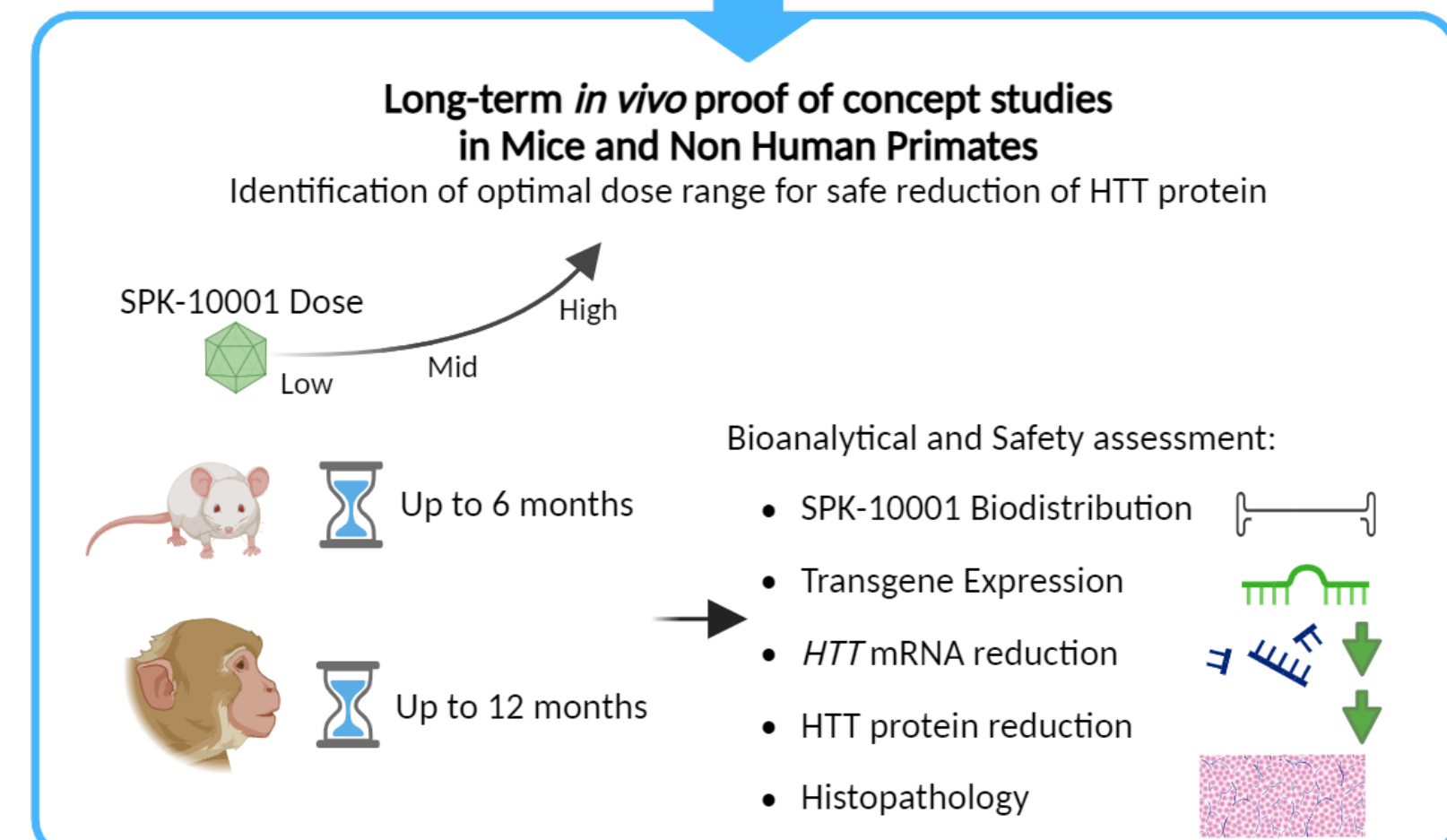
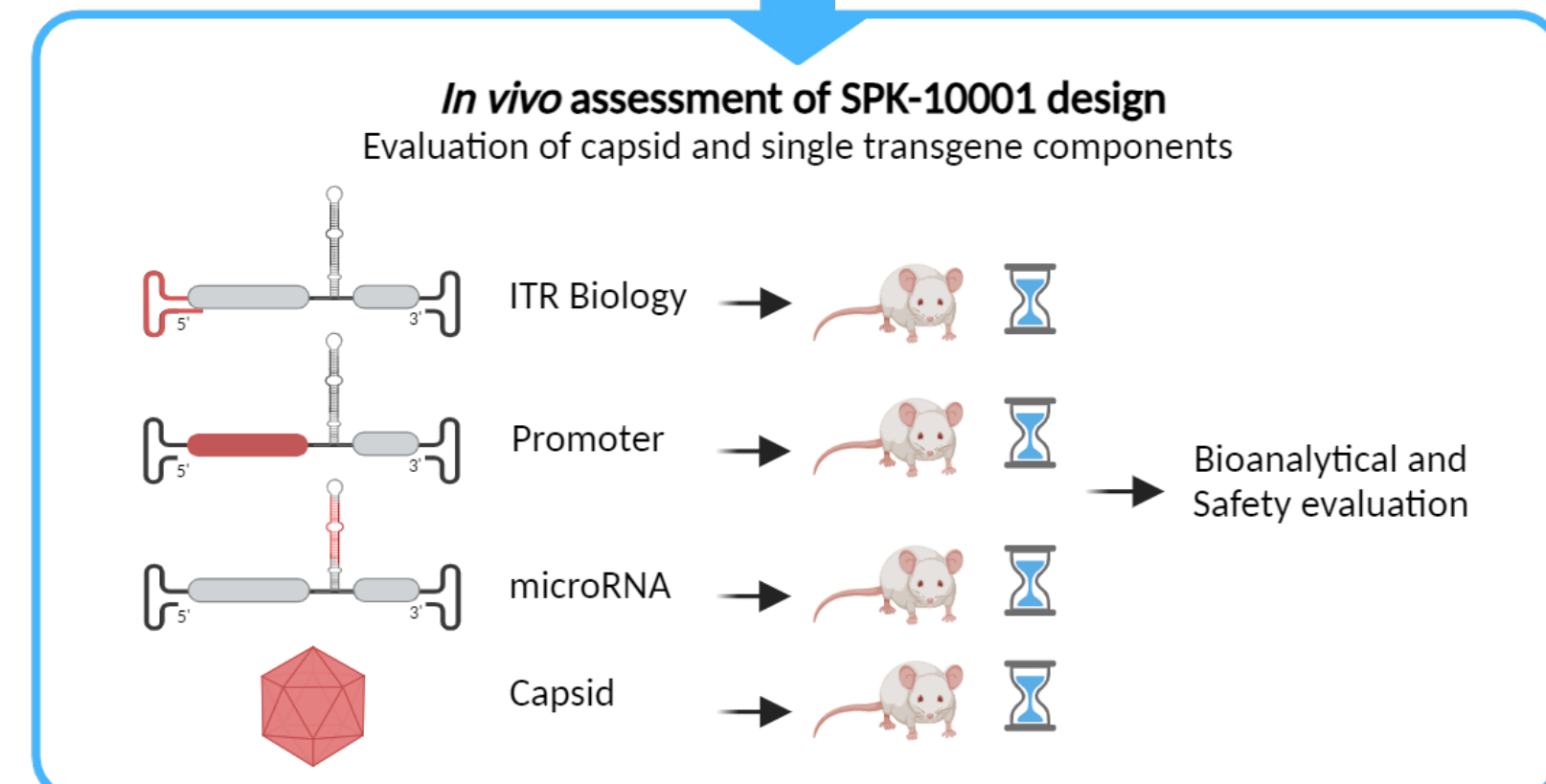
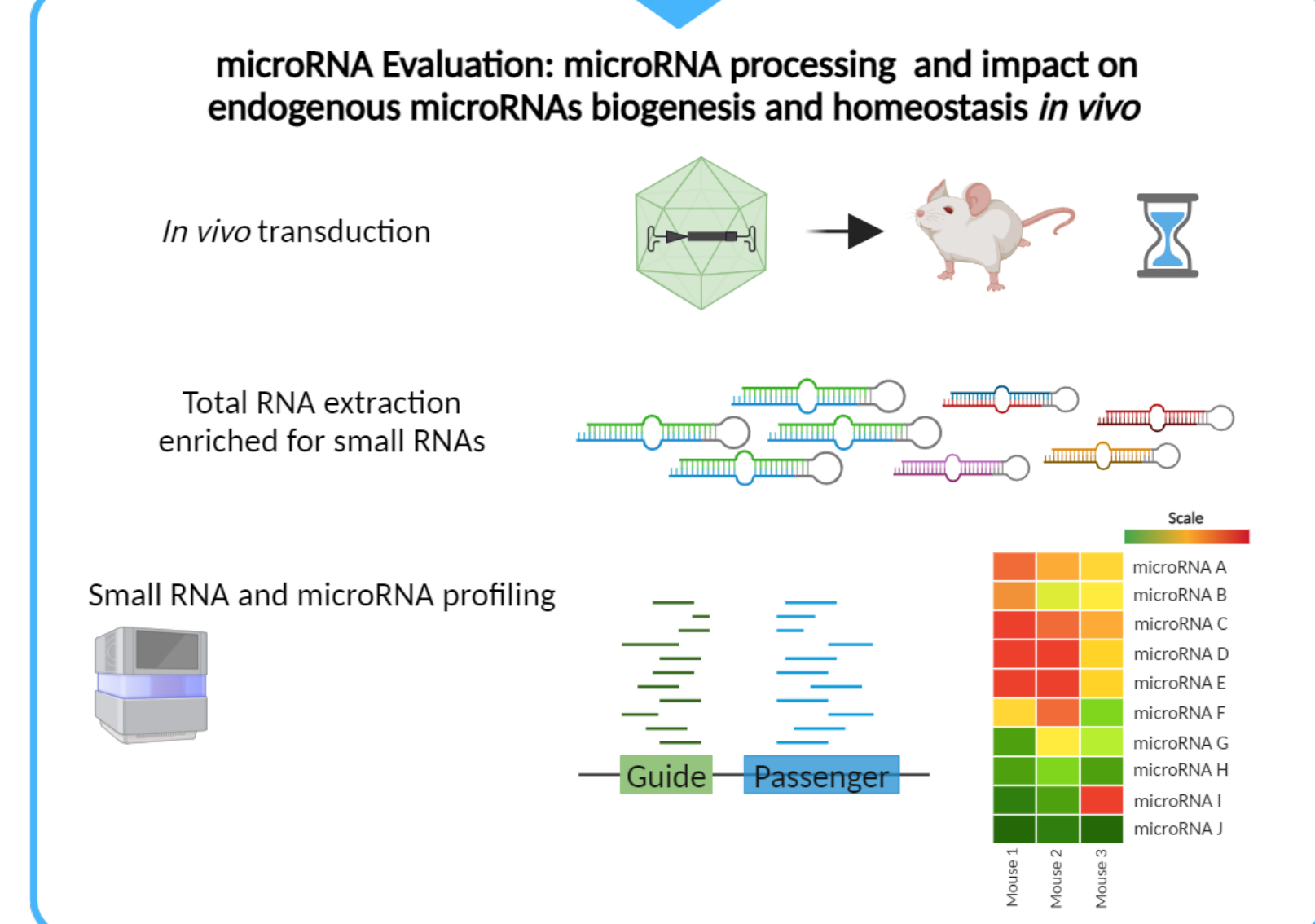
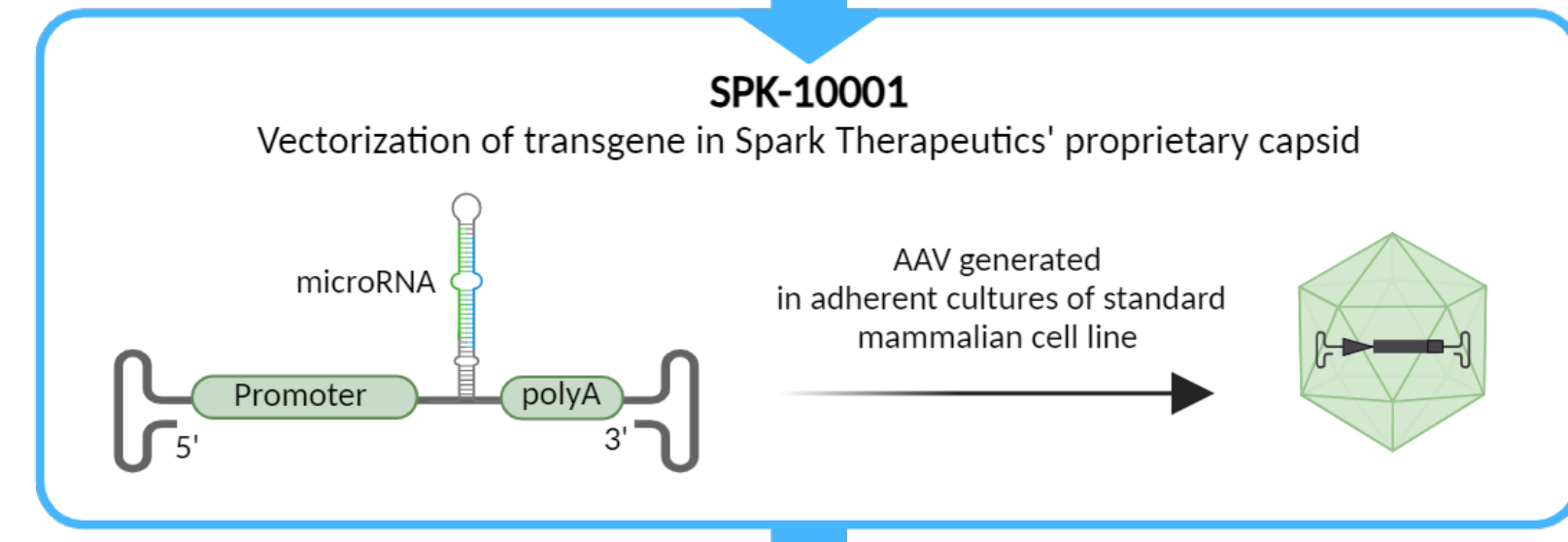
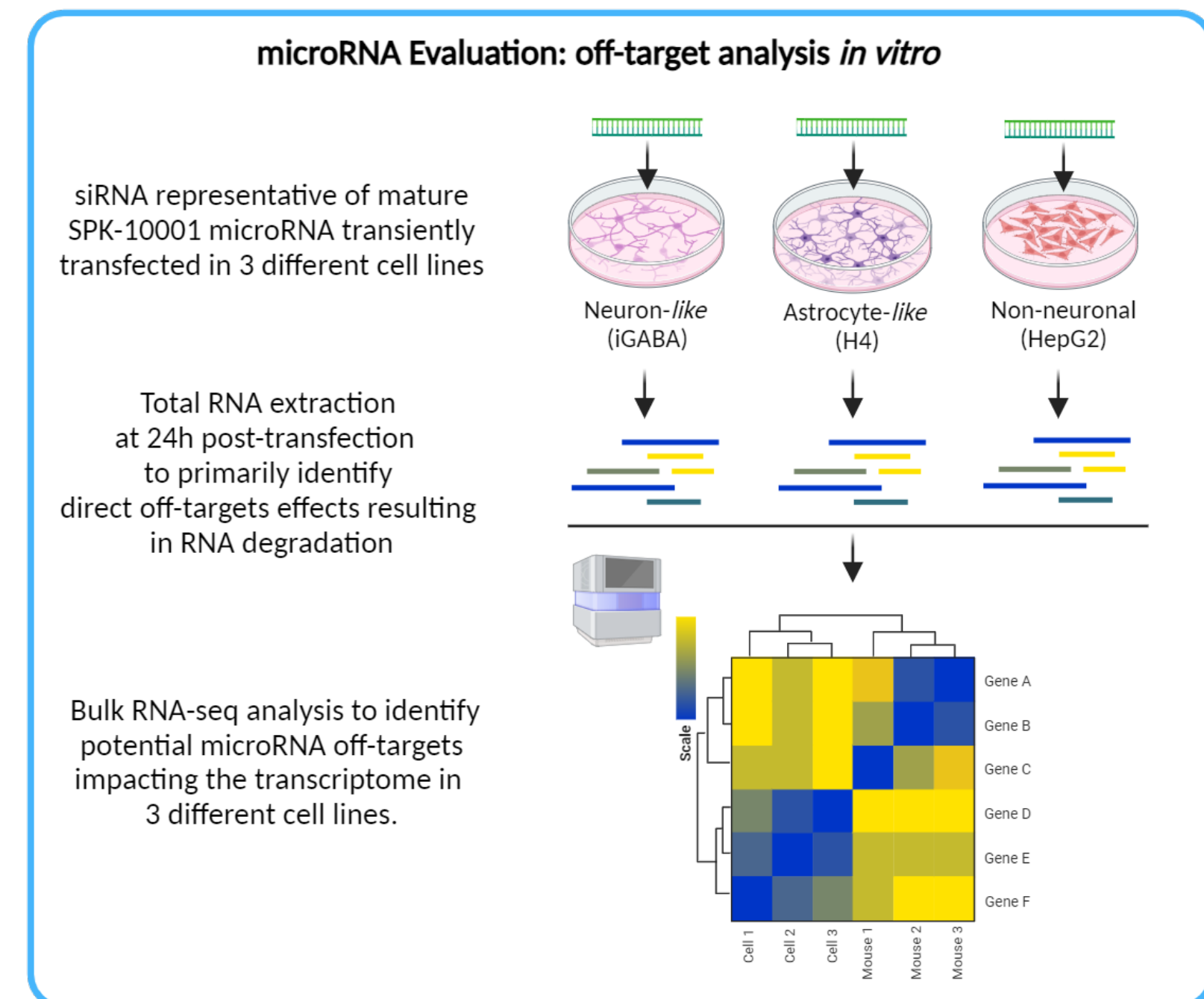
Rationale:

Huntington's disease (HD) is a fatal neurodegenerative disorder for which there are currently no disease-modifying therapies. Clinical symptoms of HD are caused by the accumulation of misfolded and aggregation-prone mutant huntingtin protein (mHTT). Reducing the amount of mHTT in neurons should in theory slow or halt the progression of disease, especially when the therapy is initiated before neurodegeneration is advanced. Advancement of mHTT-suppressing therapies has been impeded in part because many therapeutics do not efficiently penetrate the deep brain structures initially affected by disease pathology. AAV-based vectors circumvent this problem when directly administered to affected brain regions and can potentially suppress mHTT via inclusion of HTT-targeting miRNA, zinc finger proteins or similar cargo.

Approach:

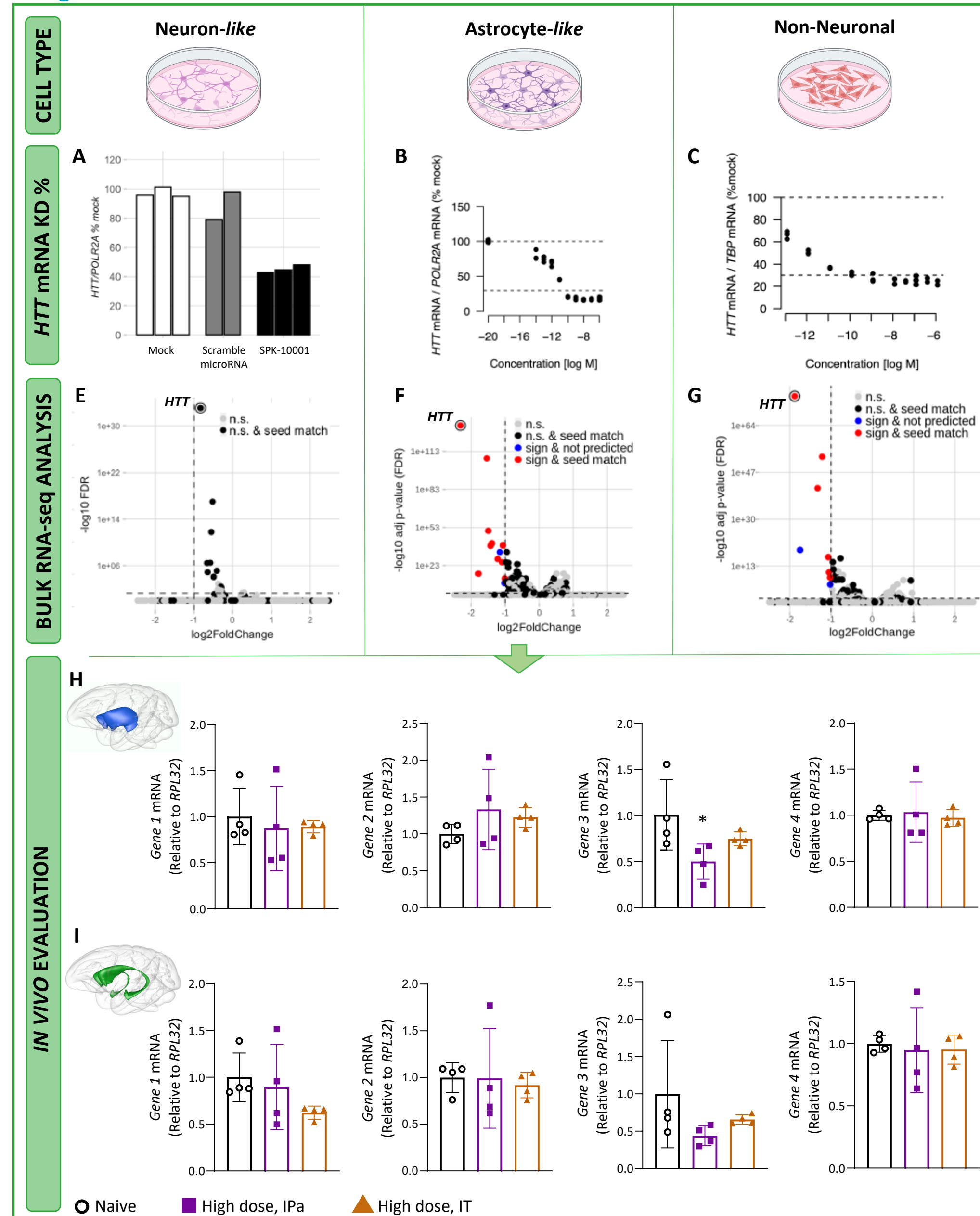
SPK-10001 comprises an engineered miRNA (HTT-miR) which binds to human *HTT* mRNA with 100% complementarity and targets it for degradation. HTT-miR expression is under control of a ubiquitous promoter and the transgene is vectorized in a proprietary AAV capsid. The HTT-miR also has 100% complementarity to macaque *HTT* mRNA which enables potency testing of SPK-10001 in non-human primates. SPK-10001 is designed to be delivered by direct injection to the caudate and putamen thereby providing durable and potent suppression of *HTT* mRNA and protein. The HTT-miR targets a region outside the exon 1 trinucleotide expansion, leading to the reduction of both normal and expanded HTT protein in non-clinical species or human patients in which both mRNAs are expressed.

METHODS



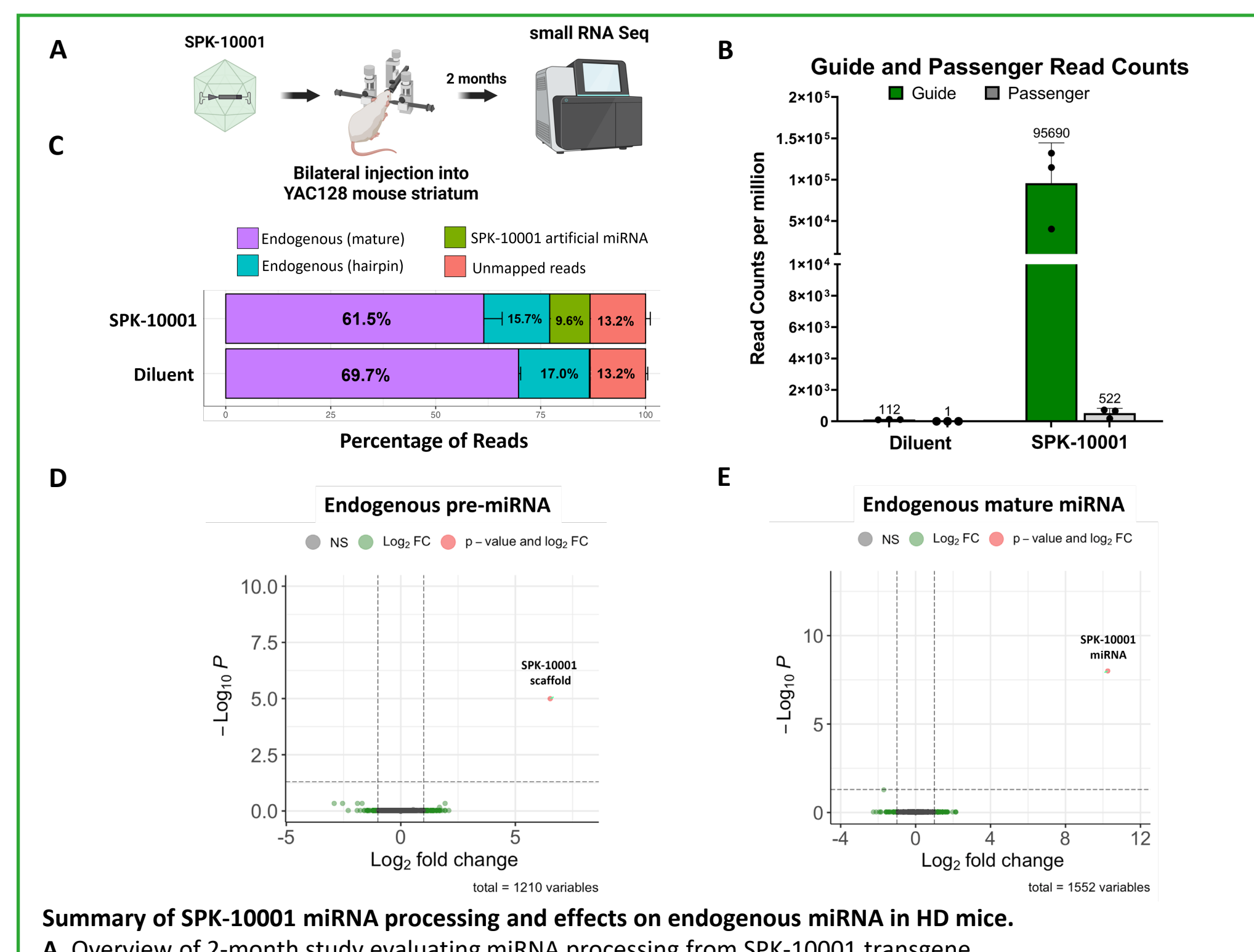
RESULTS

Figure 1. SPK-10001 microRNA has minimal effects on non-target mRNAs



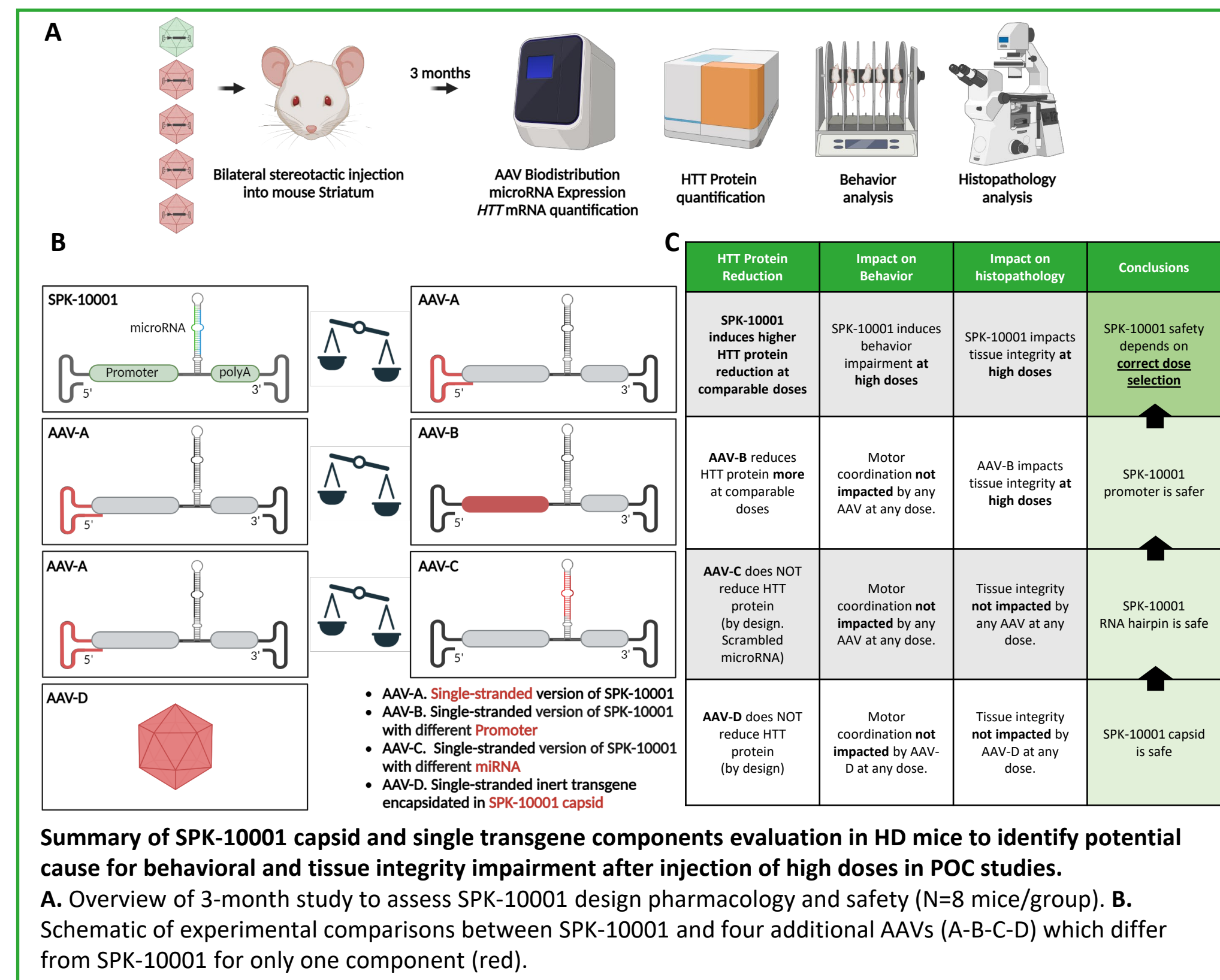
Summary of dose response curves of a siRNA version of SPK-10001 microRNA *in vitro*, identification of potential off-targets, and evaluation of potential off-targets *in vivo*.
A-B-C. Effects of siRNA targeting *HTT* mRNA as measured by qRT-PCR in A) IGABA cell line, B) H4 cell line, C) HepG2 cell line. siRNA was transfected at 10 different concentrations ranging from 10^{13} up to 10^6 M. Total RNA was extracted after 24h post-transfection and subjected to bulk sequencing to identify off-targets impacting the transcriptome and primarily identify direct off-target effects resulting in RNA degradation. Data on graphs in Panels A, B and C represents the average of three replicate experiments for each concentration. Dashed lines at 100% and 30% to help guide the eye.
D-F-G. Volcano plots displaying differentially regulated genes in D) IGABA cell line, E) H4 cell line, F) HepG2 cell line. Significantly de-regulated genes with adj p-value (FDR) < 0.05 and $\log_2 FC < -1$ are color-coded to denote predicted seed match (red) or not predicted match (blue). Non significantly de-regulated genes are color-coded based on predicted (black) or not predicted (grey) seed match. *HTT* is marked with a black ring.
H, I. Evaluation of four potential off-target mRNAs (*Gene 1*, *Gene 2*, *Gene 3*, and *Gene 4*) identified from the *in vitro* screening in NHPs Putamen (H) and Caudate (I) after intraparenchymal (IPa) or intrathecal (IT) injection of SPK-10001. Data plotted as individual NHP (symbols) and group mean \pm SD (bars). Statistical pairwise comparisons between naive and vector-injected NHPs were assessed by 1-way ANOVA followed by a Dunnett's test and reported as follows: * $p < 0.05$.

Figure 2. SPK-10001 microRNA *in vivo* processing and overexpression are efficient and safe



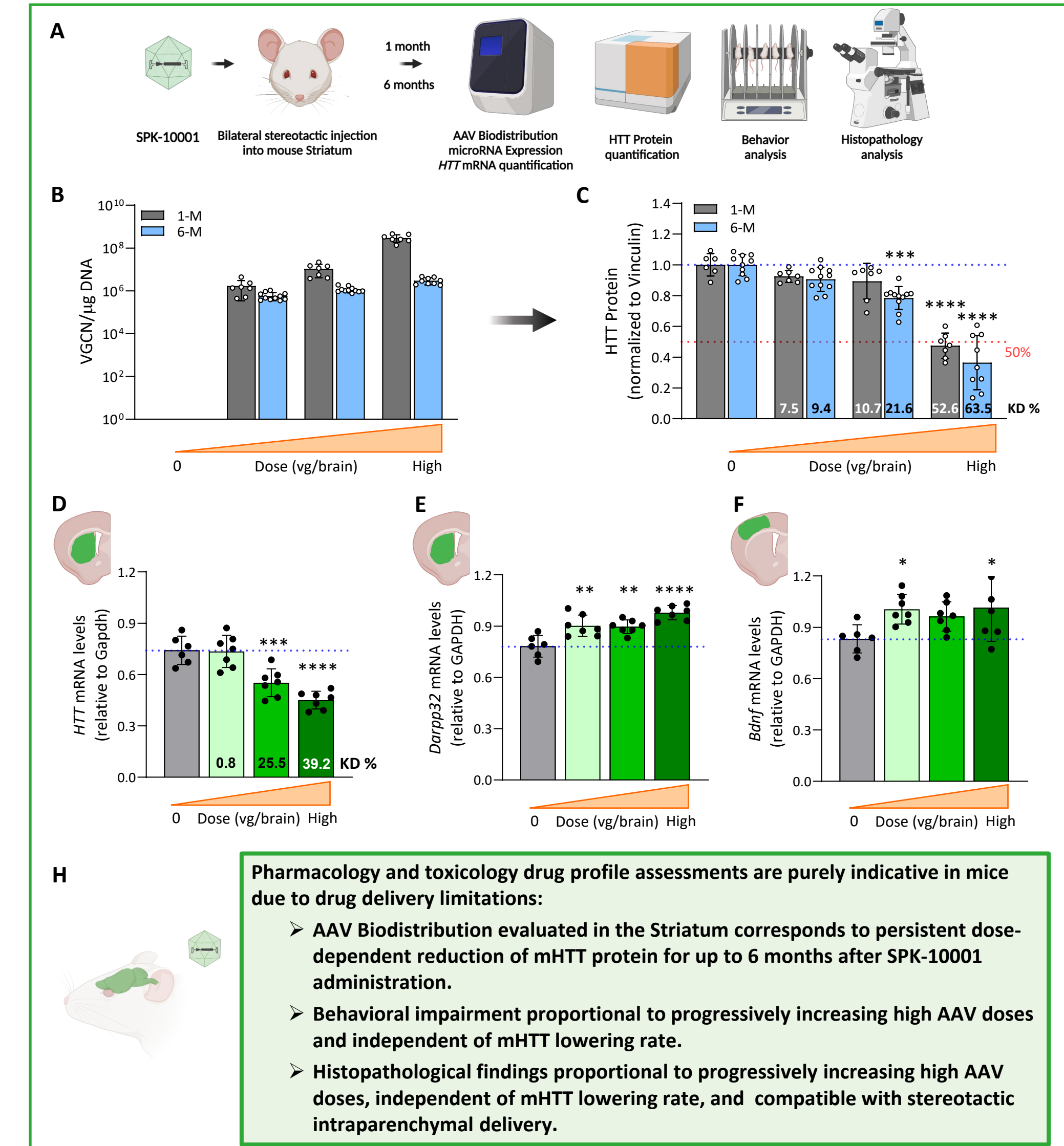
Summary of SPK-10001 miRNA processing and effects on endogenous miRNA in HD mice.
A. Overview of 2-month study evaluating miRNA processing from SPK-10001 transgene.
B. Guide and passenger read counts from YAC128 mouse striatum injected with diluent or SPK-10001. N=3 mice/group.
C. Read mapping of endogenous vs SPK-10001 artificial miRNA in YAC128 mouse striatum. N=3 mice/group.
D-E. Volcano plots showing changes in D) endogenous mouse pre-miRNA hairpins or E) endogenous mouse mature miRNAs after treatment with SPK-10001. N = 3 mice/group. No significant differences were found (adjusted $p < 0.05$ and $\log_2 FC > 1$).

Figure 3. All transgene components of SPK-10001 are well tolerated in YAC128 mice



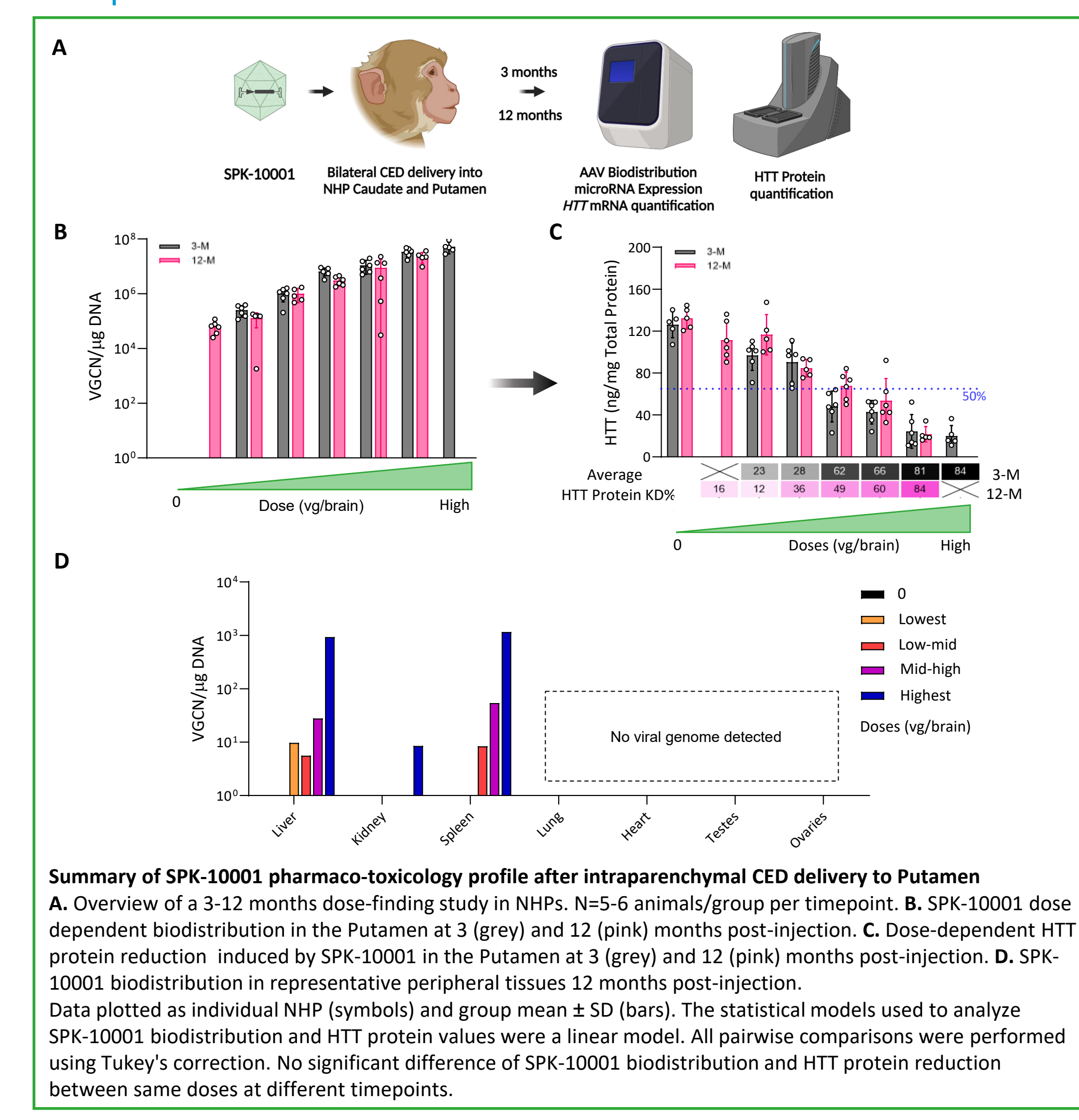
Summary of SPK-10001 capsid and single transgene components evaluation in HD mice to identify potential cause for behavioral and tissue integrity impairment after injection of high doses in POC studies.
A. Overview of 3-month study to assess SPK-10001 design pharmacology and safety (N=8 mice/group). **B.** Schematic of experimental comparisons between SPK-10001 and four additional AAVs (A-B-C-D) which differ from SPK-10001 for only one component (red). **C.** Summary of pharmacology and safety results.

Figure 4. Treatment of YAC128 mice with SPK-10001 induces durable reduction of mHTT protein and maintains the expression of *Darpp32* and *Bdnf*



Summary of SPK-10001 pharmacology-toxicology profile after intraparenchymal stereotaxic delivery to HD mice striatum.
A. Overview of a 1-6 months dose-response exploratory studies in HD mice. N=6 (1 month) -15 (6 months) animals/group. **B.** SPK-10001 dose-dependent biodistribution in the Striatum at 1 (grey) and 6 (blue) months post-injection. **C.** Dose-dependent mHTT protein reduction induced by SPK-10001 in the Striatum at 1 (grey) and 6 (blue) months post-injection. **D.** Striatal normalized *HTT* mRNA levels at 1 month after injection. **E.** Striatal normalized *Darpp32* mRNA levels at 1 month after injection. **F.** Cortical normalized *Bdnf* mRNA levels at 1 month after injection. **H.** Overall SPK-10001 pharmacology-toxicology profile as determined by preliminary dose-escalation studies.
 Data plotted as individual mouse (symbols) and group mean \pm SD (bars). Statistical pairwise comparisons between naive and vector-injected YAC128 mice were assessed by 1-way ANOVA followed by a Dunnett's test and reported as follows: * $p < 0.05$, ** $p < 0.001$, *** $p < 0.001$, **** $p < 0.0001$.

Figure 5. SPK-10001 induces an efficient and durable reduction of HTT protein in NHPs



Summary of SPK-10001 pharmacology-toxicology profile after intraparenchymal CED delivery to Putamen
A. Overview of a 3-12 months dose-finding study in NHPs. N=5-6 animals/group per timepoint. **B.** SPK-10001 dose dependent biodistribution in the Putamen at 3 (grey) and 12 (pink) months post-injection. **C.** Dose-dependent HTT protein reduction induced by SPK-10001 in the Putamen at 3 (grey) and 12 (pink) months post-injection. **D.** SPK-10001 biodistribution in representative peripheral tissues 12 months post-injection. Data plotted as individual NHP (symbols) and group mean \pm SD (bars). The statistical models used to analyze SPK-10001 biodistribution and HTT protein values were a linear model. All pairwise comparisons were performed using Tukey's correction. No significant difference of SPK-10001 biodistribution and HTT protein reduction between same doses at different timepoints.

CONCLUSIONS

- SPK-10001 is an engineered adeno-associated virus expressing an artificial microRNA which targets the human *HTT* mRNA for degradation
- SPK-10001 has minimal effects on non-target mRNAs
- microRNA overexpression induced by SPK-10001 did not affect endogenous microRNA biogenesis and homeostasis
- Persistent reduction of HTT protein up to 6 months (mice) or up to 12 months (NHPs) was well tolerated
- High doses of SPK-10001 used in pilot studies were associated with behavioral (mice) and histopathological (mice and NHPs) adverse events
- SPK-10001 will progress towards preclinical development to refine effective and safe therapeutic doses

References

1. Pfister EL, Chase KO, Sun H, Kennington LA, Conroy F, Johnson E, Miller R, Borel F, Aronin N, Mueller C. Mol Ther Nucleic Acids. 2017 Jun 16;7:324-334.

Acknowledgments

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